# Package 'PhenotypeLibraryDiagnostics'

July 12, 2022

executePhenotyeLibraryDiagnostics

Execute the cohort diagnostics

#### **Description**

Execute the cohort diagnostics

# Usage

```
executePhenotyeLibraryDiagnostics(
  connectionDetails,
  cdmDatabaseSchema,
  vocabularyDatabaseSchema = cdmDatabaseSchema,
  cohortDatabaseSchema = cdmDatabaseSchema,
  cohortTable = "cohort",
  tempEmulationSchema = getOption("sqlRenderTempEmulationSchema"),
  verifyDependencies = TRUE,
  outputFolder,
  incrementalFolder = file.path(outputFolder, "incrementalFolder"),
  databaseId = "Unknown",
  databaseName = databaseId,
  databaseDescription = databaseId
)
```

# **Arguments**

connectionDetails

An object of type connectionDetails as created using the createConnectionDetails function in the DatabaseConnector package.

cdmDatabaseSchema

Schema name where your patient-level data in OMOP CDM format resides. Note that for SQL Server, this should include both the database and schema name, for example 'cdm\_data.dbo'.

 $vocabulary {\tt Database Schema}$ 

Schema name where your OMOP vocabulary data resides. This is commonly the same as cdmDatabaseSchema. Note that for SQL Server, this should include both the database and schema name, for example 'vocabulary.dbo'.

cohortDatabaseSchema

Schema name where intermediate data can be stored. You will need to have write privileges in this schema. Note that for SQL Server, this should include both the database and schema name, for example 'cdm\_data.dbo'.

cohortTable

The name of the table that will be created in the work database schema. This table will hold the exposure and outcome cohorts used in this study.

tempEmulationSchema

Some database platforms like Oracle and Impala do not truly support temp tables. To emulate temp tables, provide a schema with write privileges where temp tables can be created.

verifyDependencies

Check whether correct package versions are installed?

uploadResults 3

 $output Folder \qquad Name \ of \ local \ folder \ to \ place \ results; \ make \ sure \ to \ use \ forward \ slashes \ (/). \ Do$ 

not use a folder on a network drive since this greatly impacts performance.

incrementalFolder

Name of local folder to hold the logs for incremental run; make sure to use forward slashes (/). Do not use a folder on a network drive since this greatly

impacts performance.

databaseId A short string for identifying the database (e.g. 'Synpuf').

databaseName The full name of the database (e.g. 'Medicare Claims Synthetic Public Use Files

(SynPUFs)').

databaseDescription

A short description (several sentences) of the database.

#### **Details**

This function executes the cohort diagnostics.

uploadResults

Upload results to OHDSI server

#### **Description**

Upload results to OHDSI server

# Usage

uploadResults(outputFolder, privateKeyFileName, userName)

## **Arguments**

outputFolder

Name of local folder where the results were generated; make sure to use forward

slashes (/).

 $\verb"privateKeyFileName"$ 

A character string denoting the path to the RSA private key provided by the

study coordinator.

userName A character string containing the user name provided by the study coordinator.

# **Details**

This function uploads the 'AllResults\_<databaseId>.zip' to the OHDSI SFTP server. Before sending, you can inspect the zip file, wich contains (zipped) CSV files. You can send the zip file from a different computer than the one on which is was created.

# Index

```
\label{lem:createConnectionDetails,2} \\ execute Phenotye Library Diagnostics,2 \\ \\ upload Results,3 \\ \\
```